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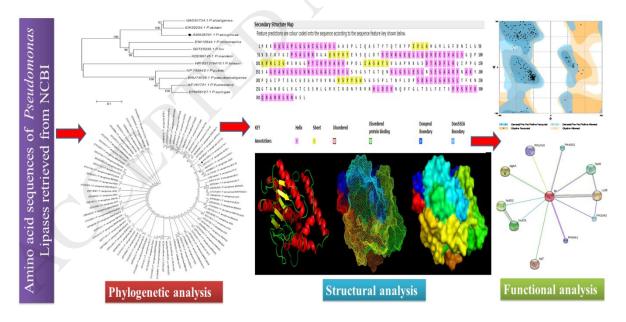
Computational elucidation of phylogenetic, structural and functional characteristics of *Pseudomonas* Lipases

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